

123-03_US_replacement_sequence.txt
SEQUENCE LISTING

<110> Hexima Limited
Poon, Simon
Heath, Robyn L.
Clarke, Adrienne E.

<120> Arabinogalactan Protein Compositions and Methods for Fostering
Somatic Embryonic Competence

<130> 12639240/AJH

<140> 10/594,418
<141> 2005-03-31

<150> 60/558,609
<151> 2004-03-01

<160> 27

<170> PatentIn version 3.4

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123-03_US_replacement_sequence.txt

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123-03_US_replacement_sequence.txt

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tcttctgaat cagattctct caacaaatgg gctgaaaaag ctcgtttcca aatcggcgac 180
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gcaccttcgc cggttgattt tgaagggtccg gccgttgctc caacaagcg agttgcaggg 480
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Lys Thr Gly Ala Trp Lys Ile Pro Ser Ser Glu Ser Asp Ser Leu Asn
35 40 45

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123-03_US_replacement_sequence.txt

Lys Trp Ala Glu Lys Ala Arg Phe Gln Ile Gly Asp Ser Leu Val Trp
50 55 60

Lys Tyr Asp Gly Gly Lys Asp Ser Val Leu Gln Val Ser Lys Glu Asp
65 70 75 80

Tyr Thr Ser Cys Asn Thr Ser Asn Pro Ile Ala Glu Tyr Lys Asp Gly
85 90 95

Asn Thr Lys Val Lys Leu Glu Lys Ser Gly Pro Tyr Phe Phe Met Ser
100 105 110

Gly Ala Lys Gly His Cys Glu Gln Gly Gln Lys Met Ile Val Val Val
115 120 125

Met Ser Gln Lys His Arg Tyr Ile Gly Ile Ser Pro Ala Pro Ser Pro
130 135 140

Val Asp Phe Glu Gly Pro Ala Val Ala Pro Thr Ser Gly Val Ala Gly
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Leu Lys Ala Gly Leu Leu Val Thr Val Gly Val Leu Gly Leu Phe
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<210> 19

<211> 660

<212> DNA

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gagaactaca atcattgggc tgaaaggaat agattccaag tcaatgatac tctctttttc	180
aagtacaaga aagggtcaga ctcggtgctg ttggttaaca gagaagatta cttctcatgc	240
aacaccaaga acccaattca gtctttaaca gaagggtgatt cactctttac atttgatcgg	300
tcgggtccct tctttttcat caccggtaac gctgataatt gcaaaaaagg gcaaaagctg	360
atcgtctggtg tcattggctgt aagacacaaa cccagcaac aacctcttc accttctccc	420
tcattctgctg tgacaacagc gccggtttct ccaccacat taccatttc tgaaactaac	480
cctcctgtag agtcacaaa gagcagttag gctccatctc atgatgctgt ggaaccagct	540
ccgcgggagc acagatcggg ttcattcaaa ctagtatgtt ctacctggct ggtgttggt	600
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123-03_US_replacement_sequence.txt

<210> 20
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Asp Gly Trp Val Val Ser Pro Ser Glu Asn Tyr Asn His Trp Ala Glu
      35      40      45

Arg Asn Arg Phe Gln Val Asn Asp Thr Leu Phe Phe Lys Tyr Lys Lys
      50      55      60

Gly Ser Asp Ser Val Leu Leu Val Thr Arg Glu Asp Tyr Phe Ser Cys
65      70      75      80

Asn Thr Lys Asn Pro Ile Gln Ser Leu Thr Glu Gly Asp Ser Leu Phe
      85      90      95

Thr Phe Asp Arg Ser Gly Pro Phe Phe Phe Ile Thr Gly Asn Ala Asp
      100      105      110

Asn Cys Lys Lys Gly Gln Lys Leu Ile Val Val Val Met Ala Val Arg
      115      120      125

His Lys Pro Gln Gln Gln Pro Pro Ser Pro Ser Pro Ser Ser Ala Val
      130      135      140

Thr Thr Ala Pro Val Ser Pro Pro Thr Leu Pro Ile Pro Glu Thr Asn
145      150      155      160

Pro Pro Val Glu Ser Pro Lys Ser Ser Glu Ala Pro Ser His Asp Ala
      165      170      175

Val Glu Pro Ala Pro Pro Glu His Arg Ser Gly Ser Phe Lys Leu Val
      180      185      190

Cys Ser Thr Trp Leu Val Leu Gly Phe Gly Ile Trp Val Ser Met Ala
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Leu Gly Ile Glu Asn Val Val Cys Phe Trp Cys
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123-03_US_replacement_sequence.txt

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123-03_US_replacement_sequence.txt

20

25

30

Gly Ser Lys Glu Ile Met Val Gly Gly Lys Thr Gly Ala Trp Lys Ile
 35 40 45

Pro Ser Ser Glu Ser Asp Ser Leu Asn Lys Trp Ala Glu Lys Ala Arg
 50 55 60

Phe Gln Ile Gly Asp Ser Leu Val Trp Lys Tyr Asp Gly Gly Lys Asp
 65 70 75 80

Ser Val Leu Gln Val Ser Lys Glu Asp Tyr Thr Ser Cys Asn Thr Ser
 85 90 95

Asn Pro Ile Ala Glu Tyr Lys Asp Gly Asn Thr Lys Val Lys Leu Glu
 100 105 110

Lys Ser Gly Pro Tyr Phe Phe Met Ser Gly Ala Lys Gly His Cys Glu
 115 120 125

Gln Gly Arg Lys Met Ile Val Val Val Met Ser Gln Lys His Arg Tyr
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Ile Gly Ile
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Gly Ser Tyr Lys Phe Tyr Val Gly Gly Arg Asp Gly Trp Val Val Ser
 35 40 45

Pro Ser Glu Asn Tyr Asn His Trp Ala Glu Arg Asn Arg Phe Gln Val
 50 55 60

Asn Asp Thr Leu Phe Phe Lys Tyr Lys Lys Gly Ser Asp Ser Val Leu
 65 70 75 80

123-03_US_replacement_sequence.txt

Leu Val Thr Arg Glu Asp Tyr Phe Ser Cys Asn Thr Lys Asn Pro Ile
85 90 95

Gln Ser Leu Thr Glu Gly Asp Ser Leu Phe Thr Phe Asp Arg Ser Gly
100 105 110

Pro Phe Phe Phe Ile Thr Gly Asn Ala Asp Asn Cys Lys Lys Gly Gln
115 120 125

Lys Leu Ile Val Val Val Met Ala Val Arg His Lys Pro Gln Gln Gln
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